

Amendments to the Claims:

This listing of claims replaces all prior versions and listings of claims in the application:

Listing of Claims:

1. (original) An isolated transcription factor that comprises at least one zinc finger domain, wherein presence of the transcription factor in a cell can alter the differentiation state of the cell.
2. (original) The transcription factor of claim 1 wherein the cell is a vertebrate cell, and presence of the transcription factor in the cell can induce a neuronal phenotype in the cell.
3. (original) The transcription factor of claim 2 wherein the cell is a vertebrate cell, and presence of the transcription factor in the cell can induce neurite extension in the cell.
4. (currently amended) The transcription factor of claim 2 wherein the DNA contacting residues of the zinc finger domain at positions -1, [[1,]] 2, 3, and 6 of the domain correspond to a motif selected from the group consisting of: QSNR, QSNK, and CSNR.
5. (currently amended) The transcription factor of claim 2 wherein the transcription factor further comprises a second and third zinc finger domain, and the DNA contacting residues of the first, second, and third domains at positions -1, [[1,]] 2, 3, and 6 of each domain respectively correspond to the motifs: QSNR, QSNK, and CSNR.
6. (original) The transcription factor of claim 5 wherein the transcription factor comprises the amino acid sequence of the zinc finger array in SEQ ID NO:2.

7. (original) The transcription factor of claim 1 wherein the cell is a vertebrate cell, and presence of the transcription factor in the cell can induce expression of an osteoblast marker in the cell.

8. (original) The transcription factor of claim 7 wherein the osteoblast marker is alkaline phosphatase.

9. (original) The transcription factor of claim 7 wherein the cell is a myoblast cell line.

10. (currently amended) The transcription factor of claim 7 wherein the DNA contacting residues of the zinc finger domain at positions -1, 1, 2, 3, and 6 of the domain correspond to a motif selected from the group consisting of: RDKR, QTHR, VSTR, and RDKR.

11. (currently amended) The transcription factor of claim 7 wherein the transcription factor further comprises a second and third zinc finger domain, and the DNA contacting residues of the first, second, and third domains at positions -1, 1, 2, 3, and 6 of each domain respectively correspond to (i) the motifs: RDKR, QTHR, and VSTR or (ii) the motifs: QTHR, VSTR, and RDKR.

12. (original) The transcription factor of claim 11 wherein the transcription factor comprises the amino acid sequence of the zinc finger array in SEQ ID NO:4.

13. (original) A nucleic acid that encodes the transcription factor of claim 1.

14. (original) A cell that contains the transcription factor of claim 1.

15. (original) A method of altering the differentiation state of a cell, the method comprising:

expressing a nucleic acid that comprises a sequence encoding the protein of claim 1 in the cell; and

maintaining the cell under conditions that allow the protein to alter the differentiated state of the cell.

16. (original) A method of altering the differentiation state of a vertebrate cell, the method comprising:

contacting the protein of claim 1 to the cell; and

maintaining the cell under conditions that allow the protein to alter the differentiated state of the cell.

17. (original) A method of altering the differentiation state of a cell in a subject, the method comprising:

administering the protein of claim 1 or a nucleic acid that comprises a sequence encoding the protein to a subject in an amount effective to alter the differentiation state of at least one cell in the subject.

18. (original) A method of inducing neurite formation in a vertebrate cell, the method comprising:

providing a vertebrate cell that contains a nucleic acid that comprises a sequence encoding an artificial transcription factor comprising three zinc finger domains, wherein expression of the artificial transcription factor induces a neuronal phenotype in the vertebrate cell; and

maintaining the vertebrate cell under conditions in which the artificial transcription factor is produced and neurite formation is induced.

19. (currently amended) The method of claim 18 wherein the DNA contacting residues of the first, second, and third domains at positions -1, 1, 2, 3, and 6 of each domain respectively correspond to the motifs: QSNR, QSNK, and CSNR.

20. (original) A method of inducing osteogenesis in a vertebrate cell, the method comprising:

providing a vertebrate cell that contains a nucleic acid that comprises a sequence encoding an artificial transcription factor comprising three zinc finger domains, wherein expression of the artificial transcription factor induces osteogenesis in the cell; and

maintaining the vertebrate cell under conditions in which the artificial transcription factor is produced and osteogenesis is induced.

21. (currently amended) The method of claim 18 wherein the DNA contacting residues of the first, second, and third domains at positions -1, 1, 2, 3, and 6 of each domain respectively correspond to (i) the motifs: RDKR, QTHR, and VSTR or (ii) the motifs: QTHR, VSTR, and RDKR.

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Serial No. : 10/669,861  
Filed : September 24, 2003  
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Attorney's Docket No.: 12279-007002

Amendments to the Drawings:

The attached replacement sheet includes changes to Fig. 25 and replaces the original sheet including Fig. 25.

In Figure 25, the sequence identifier SEQ ID NO:262 is renumbered as SEQ ID NO:64.

Attachments following last page of this Amendment:

Replacement Sheet (1 page)  
Annotated Sheet Showing Change(s) (1 page)